161

|              | I hereby certify that, on the date shown below, this correspondence is deposited with the United States Postal Service in an envelope a SEQUENCE, ASSISTANT COMMISSIONER FOR PATENTS  ☐ under 37CFR § 1.8(a), with sufficient postage as first class ☐ under 37CFR § 1.10, as "Express Mail Post Office to Address | addressed to: BOX<br>S, WASHINGTON, D.<br>mail, or | Attorney D                | PATENT Docket No. 686Q | C        |  |  |  |  |  |  |  |  |  |  |
|--------------|--|--|---------------------------|------------------------|----------|--|--|--|--|--|--|--|--|--|--|
| PE           | transmitted by facsimile to the Patent and Trademark Office, Fa  Attention: Examiner   | CN 02  | 28008                     | /                      |          |  |  |  |  |  |  |  |  |  |  |
| 4 0 3 50g    | De: November, 2001 By:Lois E. Miller   | Mew  | - #/2                     | , 1/12/<br>B           | 10       |  |  |  |  |  |  |  |  |  |  |
| TRANSM       | IN THE UNITED STATES PATENT AND TRADEMARK OFFICE   |  |                           |                        |          |  |  |  |  |  |  |  |  |  |  |
| THE STATE OF | In re application of:  | Examiner:  | Jamroz, M                 | JAN<br>CH CEN          | П        |  |  |  |  |  |  |  |  |  |  |
|              | GORMAN & MATTSON   | Art Unit:  | 1644                      | 0 9<br>TER             |          |  |  |  |  |  |  |  |  |  |  |
|              | Serial No.: 09/671,658   |  | READABLE<br>SUBMISSION    | 2002                   | <u> </u> |  |  |  |  |  |  |  |  |  |  |
|              | Filed: September 27, 2000  |  |                           | 900                    | C        |  |  |  |  |  |  |  |  |  |  |
|              | For: MAMMALIAN CELL SURFACE<br>ANTIGENS; RELATED REAGENTS  |  | alifornia 94304<br>, 2001 |                        |          |  |  |  |  |  |  |  |  |  |  |
| 5            |  | 1  |                           |                        |          |  |  |  |  |  |  |  |  |  |  |

BOX: Sequence Assistant Commissioner for Patents Washington, D.C. 20231

10 Sir:

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## COMPLIANCE WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

In reply to receipt of a "Notice to Comply" with requirements for patent applications containing nucleotide sequence and/or amino acid sequence disclosures dated November 5, 2001, for the above-identified application, in accordance with 37 CFR § 1.821 - 1.825, Applicants hereby submit: (1) a write-protected diskette containing a computer-readable submission for the "Sequence Listing"; and (2) a "Sequence Listing" paper copy of the contents of the diskette.

## REMARKS

generated by the Patent Office's PATENTIN program. The Diskette should comply with the requirements of 37 CFR §1.824 and is IBM PC compatible with a PC-DOS/MS-DOS operating system. If the diskette has been damaged, please call Applicants and a replacement diskette will be provided. A hard paper copy printout of the diskette is attached thereto.

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I hereby state the informational contents of the paper and computer readable copies of the Sequence Listing, submitted in accordance with 37 CFR 1.821(c) and (e), respectively, are believed to be the same. This submission introduces no new matter, since enclosed sequences are the same as sequences which were submitted in priority documents.

Applicants have invested significant labor and care in preparing the present submission. The enclosed items are a bona fide effort to bring the present application into full compliance with the rules for sequence submissions. Should this not be the case, Applicants respectfully request notification of specific deficiencies and an opportunity for remedy, as described in 37 CFR 1.135(c).

Applicants believe that no fees are required; however, if any fees are required by the present Response, the Commissioner is authorized to charge any fees or credit any overpayment to DNAX Research Institute Deposit Account No. 04-1239.

Respectfully submitted,

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Date: November 9, 2001

Shade Mahan

Sheela Mohan-Peterson Attorney for Applicants

Reg. No. 41,201

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enclosures and attachments:

one write-protected diskette (CRM)

paper copy of contents of diskette (4 pages)

Copy of "Notice to Comply" with sequence listing rules

DNAX Research Institute

901 California Avenue

Palo Alto, California 94304-1104 Main: (650) 496-6400

Direct:

(650) 496-1244

Fax:

(650) 496-1200

Application/Control Number: 09/671,658

Art Unit: 1644



**DETAILED ACTION** 

Sequence Compliance

1. This application contains sequence disclosures that are encompassed by the definitions for nucleotide and/or amino acid sequences set forth in 37 CFR 1.821(a)(1) and (a)(2). However, this application fails to comply with the requirements of 37 CFR 1.821 through 1.825 for the reason(s) set forth on the attached Notice To Comply With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures.

Applicant is reminded to amend the specification (including the Brief Description of Drawings) and claims as appropriate to reflect compliance with the Sequence Rules.

2. Any inquiry concerning this communication or earlier communications from the examiner should be directed to Megan Jamroz, whose telephone number is (703) 308-8365. The examiner can normally be reached Monday to Friday from 8:00 to 4:30. A message may be left on the examiner's voice mail service. If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Christina Chan can be reached at (703) 308-3973. Any inquiry of a general nature or relating to the status of this application should be directed to the Technology Center 1600 receptionist whose telephone number is (703) 308-0196.

Papers related to this application may be submitted to Technology Center 1600 by facsimile transmission. Papers should be faxed to Technology Center 1600 via the PTO Fax Center located in Crystal Mall 1. The faxing of papers must conform with the notice published in the Official Gazette, 1096 OG 30 (November 15, 1989). The CM1 Fax Center telephone number is (703) 305-3014.

Margaret (Megan) Jamroz, Ph.D. Patent Examiner Technology Center 1600 November 1, 2001

PHULIP GAMBEL, PH.D
PRIMARY EXAMINER
TEH CONCEN 1600

| ,        | 51178                    | Application No.       | Applicant(s)        |  |
|----------|--------------------------|-----------------------|---------------------|--|
| 2        | Notice to Comply         | <b>%</b> 09/671,658   | GORMAN ET AL.       |  |
| •        | Notice to Comply JAN 0 3 | 2007. Examiner        | Art Unit            |  |
|          | E                        | Margaret E Jamroz     | 1644                |  |
| NOTICE : | TO COMPLY WITH DECYLIPEN | HENRY COD DATENT ADDI | ICATIONS CONTAINING |  |

## NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Applicant must file the items indicated below within the time period set the Office action to which the Notice is attached to avoid abandonment under 35 U.S.C. § 133 (extensions of time may be obtained under the provisions of 37 CFR 1.136(a)).

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

| tor         | such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):   |
|-------------|--|
| $\boxtimes$ | 1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to the final rulemaking notice published at 55 FR 18230 (May 1, 1990), and 1.114 OG 29 (May 15, 1990). If the effective filing date is on or after July 1, 1998, see the final rulemaking notice published at 63 FR 29620 (June 1, 1998) and 1211 OG 82 (June 23, 1998). |
|             | 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).  |
| $\boxtimes$ | 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).  |
|             | 4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."  |
|             | 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must  |

unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).

☐ 6. The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing"
 as required by 37 C.F.R. 1.821(e).

7. Other:

Applicant Must Provide:

An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".

An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.

A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).

For questions regarding compliance to these requirements, please contact:

For Rules Interpretation, call (703) 308-4216

For CRF Submission Help, call (703) 308-4212

PatentIn Software Program Support

Technical Assistance......703-287-0200

To Purchase Patentin Software......703-306-2600

PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR REPLY

JAN OF THE WARD

SEQUENCE LISTING

<110> Groma Mattso <120> MAMMA

<110> Groman, Daniel M. Mattson, Jeanine D.

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gag gag atg ggc agc ggc ccc ggc gtc cca cac gag ggt ccg ctg cac Glu Glu Met Gly Ser Gly Pro Gly Val Pro His Glu Gly Pro Leu His

ccc gcg cct tct gca ccg gct ccg gcg ccg cca ccc gcc gcc tcc cgc Pro Ala Pro Ser Ala Pro Ala Pro Ala Pro Pro Pro Ala Ala Ser Arg 35 40 45

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50 55 60

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Ile Ser Glu Asp Ser Thr His Cys Phe Tyr Arg Ile Leu Arg Leu His

80 85 90 95

169

217

265

409

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| aag<br>Lys | gaa<br>Glu        | ctg<br>Leu<br>130 | caa<br>Gln | cac<br>His        | att<br>Ile | gtg<br>Val        | 999<br>Gly<br>135 | cca<br>Pro | cag<br>Gln        | cgc<br>Arg | ttc<br>Phe        | tca<br>Ser<br>140 | gga<br>Gly | gct<br>Ala        | cca<br>Pro | 553              |   |      |
|            | atg<br>Met<br>145 |                   |            |                   |            |                   |                   |            |                   |            |                   |                   |            |                   |            | 601              |   |      |
|            | gcc<br>Ala        |                   |            |                   |            |                   |                   |            |                   |            |                   |                   |            |                   |            | 649              |   |      |
|            | ggt<br>Gly        |                   |            |                   |            |                   |                   |            |                   |            |                   |                   |            |                   |            | 697              |   |      |
|            | gcc<br>Ala        |                   |            |                   |            |                   |                   |            |                   |            |                   |                   |            |                   |            | 745              | - |      |
|            | caa<br>Gln        |                   |            |                   |            |                   |                   |            |                   |            |                   |                   |            |                   |            | 793              |   |      |
| cat<br>His | gaa<br>Glu<br>225 | aca<br>Thr        | tcg<br>Ser | gga<br>Gly        | agc<br>Ser | gta<br>Val<br>230 | cct<br>Pro        | aca<br>Thr | gac<br>Asp        | tat<br>Tyr | ctt<br>Leu<br>235 | cag<br>Gln        | ctg<br>Leu | atg<br>Met        | gtg<br>Val | 841              |   |      |
|            | gtc<br>Val        |                   |            |                   |            |                   |                   |            |                   |            |                   |                   |            |                   |            | 889              | - |      |
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|            | tcc<br>Ser        |                   |            |                   |            |                   |                   |            |                   |            |                   |                   |            |                   |            | 985              |   |      |
| att<br>Ile | agc<br>Ser        | att<br>Ile<br>290 | cag<br>Gln | gtg<br>Val        | tcc<br>Ser | aac<br>Asn        | cct<br>Pro<br>295 | tcc<br>Ser | ctg<br>Leu        | ctg<br>Leu | gat<br>Asp        | ccg<br>Pro<br>300 | gat<br>Asp | caa<br>Gln        | gat<br>Asp | 1033             |   |      |
|            | acg<br>Thr<br>305 |                   |            |                   |            |                   |                   |            |                   |            |                   |                   | tga        | gact              | cat        | 1082             |   |      |
| ttc        | gtgg              | aac_a             | atta       | gcate             | gg a       | tgtc              | ctaga             | a tg       | tttg              | gaaa       | ctt               | cțta              | aaa a      | aatg              | gatgat     | . 1 <b>142</b> _ | - | <br> |
| gtc        | tatad             | cat q             | gtgt       | aaga              | ct a       | ctaa              | gaga              | c at       | ggcc              | cacg       | gtg               | tatg              | aaa (      | ctca              | cagccc     | 1202             |   | ٠    |

B,

tctctcttga gcctgtacag gttgtgtata tgtaaagtcc ataggtgatg ttagattcat 1262 ggtgattaca caacggtttt acaattttgt aatgatttcc taagaattga accagattgg 1322 gagaggtatt ccgatgctta tgaaaaactt acacgtgagc tatggaaggg ggtcacagtc 1382 totqqqtota accoptqqac atqtqccact gagaaccttg aaattaagaa gatgccatgt 1442 1502 cattgcaaag aaatgatagt gtgaagggtt aagttctttt gaattgttac attgcgctgg gacctgcaaa taagttcttt ttttctaatg aggagagaaa aatatatgta tttttatata 1562 atgtctaaag ttatatttca ggtgtaatgt tttctgtgca aagttttgta aattatattt 1622 gtgctatagt atttgattca aaatatttaa aaatgtctca ctgttgacat atttaatgtt 1682 ttaaatgtac agatgtattt aactggtgca ctttgtaatt cccctgaagg tactcgtagc 1742 taagggggca gaatactgtt tctggtgacc acatgtagtt tatttcttta ttcttttaa 1802 cttaatagag tcttcagact tgtcaaaact atgcaagcaa aataaataaa taaaaataaa 1862 1922 atqaatatct tqaataataa gtaggatgtt ggtcaccagg tgcctttcaa atttagaagc taattgactt taggagctga catagccaaa aaggatacat aataggctac tgaaaatctg 1982 2042 tcaqqaqtat ttatqcaatt attgaacagg tgtctttttt tacaagagct acaaattgta aattttgtit ctttttttc ccatagaaaa tgtactatag tttatcagcc aaaaaacaat 2102 ccacttttta atttagtgaa agttatttta ttatactgta caataaaagc attgtttctg 2162 2191 aatggcattt tttggtactt aaaaatggc

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Ile Ala Leu Phe Leu Tyr Phe Arg Ala Gln Met Asp Pro Asn Arg Ile 70 75 80

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Asn Ala Gly Leu Gln Asp Ser Thr Leu Glu Ser Glu Asp Thr Leu Pro 100 105 110

Asp Ser Cys Arg Arg Met Lys Gln Ala Phe Gln Gly Ala Val Gln Lys 115 120 125

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Ala Gln Pro Phe Ala His Leu Thr Ile Asn Ala Ala Ser Ile Pro Ser 165 170 175

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Gln Asp Gly Phe Tyr Tyr Leu Tyr Ala Asn Ile Cys Phe Arg His His 210 215 220

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Val Val Lys Thr Ser Ile Lys Ile Pro Ser Ser His Asn Leu Met Lys 245 250 255

Gly Gly Ser Thr Lys Asn Trp Ser Gly Asn Ser Glu Phe His Phe Tyr 260 265 270

Ser Ile Asn Val Gly Gly Phe Phe Lys Leu Arg Ala Gly Glu Glu Ile 275 280 285

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